

Rob J Toonen

List of Publications by Year in descending order

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Version: 2024-02-01

225
papers

12,368
citations

36691

53
h-index

37326

100
g-index

242
all docs

242
docs citations

242
times ranked

12833
citing authors

#	ARTICLE	IF	CITATIONS
1	Sponging up diversity: Evaluating metabarcoding performance for a taxonomically challenging phylum within a complex cryptobenthic community. <i>Environmental DNA</i> , 2022, 4, 239-253.	3.1	10
2	Unveiling hidden sponge biodiversity within the Hawaiian reef cryptofauna. <i>Coral Reefs</i> , 2022, 41, 727-742.	0.9	16
3	Metabarcoding as a tool to examine cryptic algae in the diets of two common grazing surgeonfishes, <i>Acanthurus</i> and <i>Triostegus</i> and <i>Acanthurus nigrofasciatus</i> . <i>Environmental DNA</i> , 2022, 4, 135-146.	3.1	8
4	A phylogenomic examination of Palmyra Atoll's corallimorpharian invader. <i>Coral Reefs</i> , 2022, 41, 673-685.	0.9	5
5	Quantifying the diet diversity of herbivorous coral reef fishes using systematic review and DNA metabarcoding. <i>Environmental DNA</i> , 2022, 4, 191-205.	3.1	13
6	Physiological acclimatization in Hawaiian corals following a 22-month shift in baseline seawater temperature and pH. <i>Scientific Reports</i> , 2022, 12, 3712.	1.6	9
7	Growth and survival among Hawaiian corals outplanted from tanks to an ocean nursery are driven by individual genotype and species differences rather than preconditioning to thermal stress. <i>PeerJ</i> , 2022, 10, e13112.	0.9	8
8	Coral bleaching responses to climate change across biological scales. <i>Global Change Biology</i> , 2022, 28, 4229-4250.	4.2	44
9	Assessing the vulnerability of marine life to climate change in the Pacific Islands region. <i>PLoS ONE</i> , 2022, 17, e0270930.	1.1	6
10	Increasing comparability among coral bleaching experiments. <i>Ecological Applications</i> , 2021, 31, e02262.	1.8	68
11	Evolutionary genomics of endangered Hawaiian tree snails (Achatinellidae: Achatinellinae) for conservation of adaptive capacity. <i>PeerJ</i> , 2021, 9, e10993.	0.9	7
12	Isotopic approaches to estimating the contribution of heterotrophic sources to Hawaiian corals. <i>Limnology and Oceanography</i> , 2021, 66, 2393-2407.	1.6	21
13	Environmental gradients drive physiological variation in Hawaiian corals. <i>Coral Reefs</i> , 2021, 40, 1505-1523.	0.9	8
14	Effect of species, provenance, and coral physiology on the composition of Hawaiian coral-associated microbial communities. <i>Coral Reefs</i> , 2021, 40, 1537-1548.	0.9	4
15	Dongsha Atoll is an important stepping-stone that promotes regional genetic connectivity in the South China Sea. <i>PeerJ</i> , 2021, 9, e12063.	0.9	3
16	Genomic divergence and differential gene expression between crustacean ecotypes across a marine thermal gradient. <i>Marine Genomics</i> , 2021, 58, 100847.	0.4	1
17	Poor data stewardship will hinder global genetic diversity surveillance. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	31
18	Inclusivity is key to progressing coral biodiversity research: Reply to comment by Bonito et al. (2021). <i>Molecular Phylogenetics and Evolution</i> , 2021, 162, 107135.	1.2	1

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19	Biodiversity of coral reef cryptobiota shuffles but does not decline under the combined stressors of ocean warming and acidification. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	21
20	A codeveloped management tool to determine harvest limits of introduced mud crabs,. <i>Pacific Conservation Biology</i> , 2021, 27, 418-431.	0.5	2
21	Community similarity and species overlap between habitats provide insight into the deep reef refuge hypothesis. <i>Scientific Reports</i> , 2021, 11, 23787.	1.6	1
22	CoralCam: A flexible, low-cost ecological monitoring platform. <i>HardwareX</i> , 2020, 7, e00089.	1.1	9
23	Building a global genomics observatory: Using GEOME (the Genomic Observatories Metadatabase) to expedite and improve deposition and retrieval of genetic data and metadata for biodiversity research. <i>Molecular Ecology Resources</i> , 2020, 20, 1458-1469.	2.2	32
24	The legacy of stress: Coral bleaching impacts reproduction years later. <i>Functional Ecology</i> , 2020, 34, 2315-2325.	1.7	46
25	Host-symbiont coevolution, cryptic structure, and bleaching susceptibility, in a coral species complex (<i>Scleractinia</i> ; <i>Poritidae</i>). <i>Scientific Reports</i> , 2020, 10, 16995.	1.6	33
26	Multi-model seascape genomics identifies distinct environmental drivers of selection among sympatric marine species. <i>BMC Evolutionary Biology</i> , 2020, 20, 121.	3.2	11
27	Is post-bleaching recovery of <i>Acropora hyacinthus</i> on Palau via spread of local kin groups?. <i>Coral Reefs</i> , 2020, 39, 687-699.	0.9	4
28	Evolutionary biogeography of the reef-building coral genus <i>Galaxea</i> across the Indo-Pacific ocean. <i>Molecular Phylogenetics and Evolution</i> , 2020, 151, 106905.	1.2	20
29	Species Radiations in the Sea: What the Flock?. <i>Journal of Heredity</i> , 2020, 111, 70-83.	1.0	20
30	Abundance, size, and survival of recruits of the reef coral <i>Pocillopora acuta</i> under ocean warming and acidification. <i>PLoS ONE</i> , 2020, 15, e0228168.	1.1	29
31	Genomics versus mtDNA for resolving stock structure in the silky shark (<i>Carcharhinus</i>) Tj ETQq1 1 0.784314 rgBT/Overlock 10 Tf 5 0.9 24	0.9	24
32	Unexpectedly high genetic diversity in a rare and endangered seabird in the Hawaiian Archipelago. <i>PeerJ</i> , 2020, 8, e8463.	0.9	5
33	Genetic structure is stronger across human-impacted habitats than among islands in the coral <i>Porites lobata</i> . <i>PeerJ</i> , 2020, 8, e8550.	0.9	17
34	<i>Atlantia</i> , a new genus of <i>Dendrophylliidae</i> (Cnidaria, Anthozoa, Scleractinia) from the eastern Atlantic. <i>PeerJ</i> , 2020, 8, e8633.	0.9	8
35	Title is missing!. , 2020, 15, e0228168.		0
36	Title is missing!. , 2020, 15, e0228168.		0

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37	Title is missing!. , 2020, 15, e0228168.		0
38	Title is missing!. , 2020, 15, e0228168.		0
39	The complete mitochondrial genome of the Band-rumped Storm Petrel (<i>Oceanodroma castro</i>). Mitochondrial DNA Part B: Resources, 2019, 4, 1271-1272.	0.2	3
40	Re-description of type material of <i>Xenia</i> Lamarck, 1816 (Octocorallia: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 622 T	0.2	6
41	Rare coral under the genomic microscope: timing and relationships among Hawaiian <i>Montipora</i> . BMC Evolutionary Biology, 2019, 19, 153.	3.2	16
42	RADseq population genomics confirms divergence across closely related species in blue coral (<i>Heliopora coerulea</i>). BMC Evolutionary Biology, 2019, 19, 187.	3.2	12
43	Present-Day Distribution and Potential Spread of the Invasive Green Alga <i>Avrainvillea amadelpha</i> Around the Main Hawaiian Islands. Frontiers in Marine Science, 2019, 6, .	1.2	8
44	Diversity and Structure of Parrotfish Assemblages across the Northern Great Barrier Reef. Diversity, 2019, 11, 14.	0.7	19
45	American Samoa. Coral Reefs of the World, 2019, , 387-407.	0.3	3
46	The Hawaiian Archipelago. Coral Reefs of the World, 2019, , 445-464.	0.3	11
47	Adaptive responses and local stressor mitigation drive coral resilience in warmer, more acidic oceans. Proceedings of the Royal Society B: Biological Sciences, 2019, 286, 20190614.	1.2	47
48	The molecular biogeography of the Indo-Pacific: Testing hypotheses with multispecies genetic patterns. Global Ecology and Biogeography, 2019, 28, 943-960.	2.7	43
49	Evidence of local adaptation in a waterfall-climbing Hawaiian goby fish derived from coupled biophysical modeling of larval dispersal and post-settlement selection. BMC Evolutionary Biology, 2019, 19, 88.	3.2	9
50	High heritability of coral calcification rates and evolutionary potential under ocean acidification. Scientific Reports, 2019, 9, 20419.	1.6	29
51	A coalescent sampler successfully detects biologically meaningful population structure overlooked by F _{st} statistics. Evolutionary Applications, 2019, 12, 255-265.	1.5	15
52	Annotated checklist for stony corals of American Samoa with reference to mesophotic depth records. ZooKeys, 2019, 849, 1-170.	0.5	5
53	Molecular and morphological congruence of three new cryptic <i>Neopetrosia</i> spp. in the Caribbean. PeerJ, 2019, 7, e6371.	0.9	7
54	Shared genomic outliers across two divergent population clusters of a highly threatened seagrass. PeerJ, 2019, 7, e6806.	0.9	29

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55	A behavioral and genetic study of multiple paternity in a polygamous marine invertebrate, <i>Octopus oliveri</i> . PeerJ, 2019, 7, e6927.	0.9	8
56	Hawaiian green turtles graze on bioeroding sponges at Maunalua Bay, Oâ€ˆahu, Hawaiâ€ˆi. Galaxea, 2019, 21, 3-4.	0.2	2
57	Critical Information Gaps Impeding Understanding of the Role of Larval Connectivity Among Coral Reef Islands in an Era of Global Change. Frontiers in Marine Science, 2018, 5, .	1.2	18
58	A comparison of mitochondrial genomes from five species in three genera suggests polyphyly in the subfamily Achatinellinae (Gastropoda: Pulmonata: Stylommatophora: Achatinellidae). Mitochondrial DNA Part B: Resources, 2018, 3, 611-612.	0.2	5
59	High-frequency temperature variability mirrors fixed differences in thermal limits of the massive coral <i>Porites lobata</i> (Dana, 1846). Journal of Experimental Biology, 2018, 221, .	0.8	36
60	Divergence times in demosponges (Porifera): first insights from new mitogenomes and the inclusion of fossils in a birth-death clock model. BMC Evolutionary Biology, 2018, 18, 114.	3.2	49
61	Complex signatures of genomic variation of two non-model marine species in a homogeneous environment. BMC Genomics, 2018, 19, 347.	1.2	21
62	The first Hawaiâ€ˆi workshop for coral restoration & nurseries. Marine Policy, 2018, 96, 133-135.	1.5	4
63	A linked land-sea modeling framework to inform ridge-to-reef management in high oceanic islands. PLoS ONE, 2018, 13, e0193230.	1.1	47
64	A simple molecular technique for distinguishing species reveals frequent misidentification of Hawaiian corals in the genus <i>Pocillopora</i> . PeerJ, 2018, 6, e4355.	0.9	34
65	The little shrimp that could: phylogeography of the circumtropical <i>Stenopus hispidus</i> (Crustacea: Decapoda), reveals divergent Atlantic and Pacific lineages. PeerJ, 2018, 6, e4409.	0.9	11
66	Geopolitical species revisited: genomic and morphological data indicate that the roundtail chub <i>Gila robusta</i> species complex (Teleostei, Cyprinidae) is a single species. PeerJ, 2018, 6, e5605.	0.9	8
67	Modeled larval connectivity of a multi-species reef fish and invertebrate assemblage off the coast of Molokaâ€ˆi, Hawaiâ€ˆi. PeerJ, 2018, 6, e5688.	0.9	5
68	Coral reef grazer-benthos dynamics complicated by invasive algae in a small marine reserve. Scientific Reports, 2017, 7, 43819.	1.6	26
69	Species boundaries in the absence of morphological, ecological or geographical differentiation in the Red Sea octocoral genus <i>Ovabunda</i> (Alcyonacea: Xeniidae). Molecular Phylogenetics and Evolution, 2017, 112, 174-184.	1.2	53
70	Lifeâ€ˆhistory predicts past and present population connectivity in two sympatric sea stars. Ecology and Evolution, 2017, 7, 3916-3930.	0.8	17
71	Coral hybridization or phenotypic variation? Genomic data reveal gene flow between <i>Porites lobata</i> and <i>P. Compressa</i> . Molecular Phylogenetics and Evolution, 2017, 111, 132-148.	1.2	59
72	Connecting Palauâ€™s marine protected areas: a population genetic approach to conservation. Coral Reefs, 2017, 36, 735-748.	0.9	12

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73	Scaling Up Restoration Efforts in the Pacific Islands: A Call for Clear Management Objectives, Targeted Research to Minimize Uncertainty, and Innovative Solutions to a Wicked Problem. <i>Pacific Science</i> , 2017, 71, 391-399.	0.2	7
74	Disentangling the relative merits and disadvantages of parentage analysis and assignment tests for inferring population connectivity. <i>ICES Journal of Marine Science</i> , 2017, 74, 1749-1762.	1.2	24
75	A genomic glance through the fog of plasticity and diversification in <i>Pocillopora</i> . <i>Scientific Reports</i> , 2017, 7, 5991.	1.6	87
76	Plasticity or chimerism? Color polymorphism in <i>Montipora verrilli</i> and <i>M. patula</i> . <i>Galaxea</i> , 2017, 19, 33-34.	0.2	2
77	Coral calcification mechanisms facilitate adaptive responses to ocean acidification. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2017, 284, 20172117.	1.2	70
78	The Genomic Observatories Metadatabase (GeOMe): A new repository for field and sampling event metadata associated with genetic samples. <i>PLoS Biology</i> , 2017, 15, e2002925.	2.6	72
79	Clone wars: asexual reproduction dominates in the invasive range of <i>Tubastraea</i> spp. (Anthozoa). <i>Trends in Ecology and Evolution</i> , 2017, 32, 101-109.	0.9	37
80	The implementation of rare events logistic regression to predict the distribution of mesophotic hard corals across the main Hawaiian Islands. <i>PeerJ</i> , 2016, 4, e2189.	0.9	9
81	Modeled Population Connectivity across the Hawaiian Archipelago. <i>PLoS ONE</i> , 2016, 11, e0167626.	1.1	27
82	Temporal and spatial trends in prey composition of wahoo <i>Acanthocybium solandri</i> : a diet analysis from the central North Pacific Ocean using visual and DNA barcoding techniques. <i>Journal of Fish Biology</i> , 2016, 88, 1501-1523.	0.7	14
83	Testing dispersal limits in the sea: range-wide phylogeography of the pronghorn spiny lobster <i>Panulirus penicillatus</i> . <i>Journal of Biogeography</i> , 2016, 43, 1032-1044.	1.4	32
84	The complete mitochondrial genome of the lobe coral <i>Porites lobata</i> (Anthozoa: Scleractinia) sequenced using ezRAD. <i>Mitochondrial DNA Part B: Resources</i> , 2016, 1, 247-249.	0.2	16
85	The DNA of coral reef biodiversity: predicting and protecting genetic diversity of reef assemblages. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2016, 283, 20160354.	1.2	45
86	Comparative phylogeography of the ocean planet. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 7962-7969.	3.3	190
87	The complete mitochondrial genome of <i>Achatinella mustelina</i> (Gastropoda: Pulmonata). <i>PLoS ONE</i> , 2016, 11, e0167626.	0.2	8
88	The complete mitochondrial genome of <i>Achatinella sowerbyana</i> (Gastropoda: Pulmonata). <i>PLoS ONE</i> , 2016, 11, e0167626.	0.2	6
89	Ecological and genetic variation in reef-building corals on four Society Islands. <i>Limnology and Oceanography</i> , 2016, 61, 543-557.	1.6	27
90	On the origin of endemic species in the Red Sea. <i>Journal of Biogeography</i> , 2016, 43, 13-30.	1.4	133

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91	A review of contemporary patterns of endemism for shallow water reef fauna in the Red Sea. <i>Journal of Biogeography</i> , 2016, 43, 423-439.	1.4	150
92	An assessment of shallow and mesophotic reef brachyuran crab assemblages on the south shore of Oâ€šahu, Hawaiâ€š. <i>Coral Reefs</i> , 2016, 35, 103-112.	0.9	31
93	A decade of seascape genetics: contributions to basic and applied marine connectivity. <i>Marine Ecology - Progress Series</i> , 2016, 554, 1-19.	0.9	229
94	Population genetic structure between Yap and Palau for the coral <i>Acropora hyacinthus</i> . <i>PeerJ</i> , 2016, 4, e2330.	0.9	10
95	Growing coral larger and faster: micro-colony-fusion as a strategy for accelerating coral cover. <i>PeerJ</i> , 2015, 3, e1313.	0.9	94
96	Clues to unraveling the coral species problem: distinguishing species from geographic variation in <i>Porites</i> across the Pacific with molecular markers and microskeletal traits. <i>PeerJ</i> , 2015, 3, e751.	0.9	43
97	The unnatural history of Kâ€šneâ€šohe Bay: coral reef resilience in the face of centuries of anthropogenic impacts. <i>PeerJ</i> , 2015, 3, e950.	0.9	112
98	Could polyp pulsation be the key to species boundaries in the genus <i>Ovabunda</i> (Octocorallia: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 462	1.0	9
99	Demystifying computer science for molecular ecologists. <i>Molecular Ecology</i> , 2015, 24, 2619-2640.	2.0	6
100	Cryptic species obscure introduction pathway of the blue Caribbean sponge (<i>Haliclona</i> (<i>Soestella</i>) <i>caerulea</i>), (order: Haplosclerida) to Palmyra Atoll, Central Pacific. <i>PeerJ</i> , 2015, 3, e1170.	0.9	11
101	Survivorship and feeding preferences among size classes of outplanted sea urchins, <i>Tripneustes gratilla</i> , and possible use as biocontrol for invasive alien algae. <i>PeerJ</i> , 2015, 3, e1235.	0.9	38
102	First description of hatchlings and eggs of <i>Octopus oliveri</i> (Berry, 1914) (Cephalopoda: Octopodidae). <i>Molluscan Research</i> , 2014, 34, 79-83.	0.2	5
103	A revision of the octocoral genus <i>Ovabunda</i> (Alderslade, 2001) (Anthozoa, Octocorallia, Xeniidae). <i>ZooKeys</i> , 2014, 373, 1-41.	0.5	18
104	The application of genetics to marine management and conservation: examples from the Indo-Pacific. <i>Bulletin of Marine Science</i> , 2014, 90, 123-158.	0.4	78
105	The scope of published population genetic data for Indo-Pacific marine fauna and future research opportunities in the region. <i>Bulletin of Marine Science</i> , 2014, 90, 47-78.	0.4	44
106	Comparative population structure of two edible Indo-Pacific coral reef sea cucumbers (Echinodermata: Holothuroidea). <i>Bulletin of Marine Science</i> , 2014, 90, 359-378.	0.4	10
107	Discordant population expansions in four species of coralâ€šassociated Pacific hermit crabs (Anomura: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 462	1.4	7
108	The founding charter of the Genomic Observatories Network. <i>GigaScience</i> , 2014, 3, 2.	3.3	51

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109	Archaeological Evidence of Validity of Fish Populations on Unexploited Reefs as Proxy Targets for Modern Populations. <i>Conservation Biology</i> , 2014, 28, 1322-1330.	2.4	7
110	Phylogeography unplugged: comparative surveys in the genomic era. <i>Bulletin of Marine Science</i> , 2014, 90, 13-46.	0.4	86
111	Large-scale introduction of the Indo-Pacific damselfish <i>Abudefduf vaigiensis</i> into Hawaii promotes genetic swamping of the endemic congener <i>A. abdominalis</i> . <i>Molecular Ecology</i> , 2014, 23, 5552-5565.	2.0	49
112	Demystifying the RAD fad. <i>Molecular Ecology</i> , 2014, 23, 5937-5942.	2.0	199
113	Emergent patterns of population genetic structure for a coral reef community. <i>Molecular Ecology</i> , 2014, 23, 3064-3079.	2.0	94
114	Extreme phenotypic polymorphism in the coral genus <i>Pocillopora</i> ; micro-morphology corresponds to mitochondrial groups, while colony morphology does not. <i>Bulletin of Marine Science</i> , 2014, 90, 211-231.	0.4	52
115	Regional population structure of <i>Montipora capitata</i> across the Hawaiian Archipelago. <i>Bulletin of Marine Science</i> , 2014, 90, 257-275.	0.4	32
116	After the gold rush: population structure of spiny lobsters in Hawaii following a fishery closure and the implications for contemporary spatial management. <i>Bulletin of Marine Science</i> , 2014, 90, 331-357.	0.4	9
117	Evolving coral reef conservation with genetic information. <i>Bulletin of Marine Science</i> , 2014, 90, 159-185.	0.4	89
118	Intraspecific fluorescent phenotypes in <i>Montipora capitata</i> . <i>Galaxea</i> , 2014, 16, 17-18.	0.2	1
119	A taxonomic survey of Saudi Arabian Red Sea octocorals (Cnidaria: Alcyonacea). <i>Marine Biodiversity</i> , 2013, 43, 279-291.	0.3	25
120	One size does not fit all: The emerging frontier in large-scale marine conservation. <i>Marine Pollution Bulletin</i> , 2013, 77, 7-10.	2.3	131
121	Very fine-scale population genetic structure of sympatric asterinid sea stars with benthic and pelagic larvae: influence of mating system and dispersal potential. <i>Biological Journal of the Linnean Society</i> , 2013, 108, 821-833.	0.7	31
122	The origins of tropical marine biodiversity. <i>Trends in Ecology and Evolution</i> , 2013, 28, 359-366.	4.2	377
123	The evolving male: spinner dolphin (<i>Stenella longirostris</i>) ecotypes are divergent at Y chromosome but not mtDNA or autosomal markers. <i>Molecular Ecology</i> , 2013, 22, 2408-2423.	2.0	27
124	Next-Generation Sequencing for High-Throughput Molecular Ecology: A Step-by-Step Protocol for Targeted Multilocus Genotyping by Pyrosequencing. <i>Methods in Molecular Biology</i> , 2013, 1006, 89-99.	0.4	4
125	Optimizing Selection of Microsatellite Loci from 454 Pyrosequencing via Post-sequencing Bioinformatic Analyses. <i>Methods in Molecular Biology</i> , 2013, 1006, 101-120.	0.4	3
126	Combined analyses of kinship and <i>F_{ST}</i> suggest potential drivers of chaotic genetic patchiness in high gene flow populations. <i>Molecular Ecology</i> , 2013, 22, 3476-3494.	2.0	132

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127	Population structure in the native range predicts the spread of introduced marine species. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2013, 280, 20130409.	1.2	31
128	Shallow gene pools in the high intertidal: extreme loss of genetic diversity in viviparous sea stars (<i>Tj ETQq0 0 0 rgBT/Overlock 10 Tf 50</i>)	1.0	12
129	ezRAD: a simplified method for genomic genotyping in non-model organisms. <i>PeerJ</i> , 2013, 1, e203.	0.9	184
130	Biological and Physical Interactions on a Tropical Island Coral Reef: Transport and Retention Processes on Moorea, French Polynesia. <i>Oceanography</i> , 2013, 26, 52-63.	0.5	61
131	Genetic Evidence for Regional Isolation of Pocillopora Corals from Moorea. <i>Oceanography</i> , 2013, 26, 153-155.	0.5	23
132	Microsatellites for Next-Generation Ecologists: A Post-Sequencing Bioinformatics Pipeline. <i>PLoS ONE</i> , 2013, 8, e55990.	1.1	49
133	An Invasive Fish and the Time-Lagged Spread of Its Parasite across the Hawaiian Archipelago. <i>PLoS ONE</i> , 2013, 8, e56940.	1.1	33
134	Buffer Capacity, Ecosystem Feedbacks, and Seawater Chemistry under Global Change. <i>Water (Switzerland)</i> , 2013, 5, 1303-1325.	1.2	48
135	Polyphyly and hidden species among Hawaiï™s dominant mesophotic coral genera, <i>Leptoseris</i> and <i>Pavona</i> (Scleractinia: Agariciidae). <i>PeerJ</i> , 2013, 1, e132.	0.9	43
136	Between tide and wave marks: a unifying model of physical zonation on littoral shores. <i>PeerJ</i> , 2013, 1, e154.	0.9	26
137	Growth of cultured giant clams (<i>Tridacna</i> spp.) in low pH, high-nutrient seawater: species-specific effects of substrate and supplemental feeding under acidification. <i>Journal of the Marine Biological Association of the United Kingdom</i> , 2012, 92, 731-740.	0.4	23
138	Spatial variability in growth and prey availability of lobsters in the northwestern Hawaiian Islands. <i>Marine Ecology - Progress Series</i> , 2012, 449, 211-220.	0.9	12
139	Coming out of the starting blocks: extended lag time rearranges genetic diversity in introduced marine fishes of Hawaiï™. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2012, 279, 3948-3957.	1.2	22
140	Next-Generation Phylogeography: A Targeted Approach for Multilocus Sequencing of Non-Model Organisms. <i>PLoS ONE</i> , 2012, 7, e34241.	1.1	39
141	Coral farming: effects of light, water motion and artificial foods. <i>Journal of the Marine Biological Association of the United Kingdom</i> , 2012, 92, 721-729.	0.4	22
142	The Biology and Ecology of Black Corals (Cnidaria: Anthozoa: Hexacorallia: Antipatharia). <i>Advances in Marine Biology</i> , 2012, 63, 67-132.	0.7	105
143	Molecular Delineation of Species in the Coral Holobiont. <i>Advances in Marine Biology</i> , 2012, 63, 1-65.	0.7	58
144	Extraordinarily rapid life-history divergence between Cryptasterina sea star species. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2012, 279, 3914-3922.	1.2	45

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145	Sexual reproduction of the Hawaiian black coral <i>Antipathes griggi</i> (Cnidaria: Antipatharia). <i>Coral Reefs</i> , 2012, 31, 795-806.	0.9	10
146	Global Phylogeography with Mixed-Marker Analysis Reveals Male-Mediated Dispersal in the Endangered Scalloped Hammerhead Shark (<i>Sphyrna lewini</i>). <i>PLoS ONE</i> , 2012, 7, e29986.	1.1	123
147	Sympatric Speciation in the Post "Modern Synthesis" Era of Evolutionary Biology. <i>Evolutionary Biology</i> , 2012, 39, 158-180.	0.5	89
148	Common misconceptions in molecular ecology: echoes of the modern synthesis. <i>Molecular Ecology</i> , 2012, 21, 4171-4189.	2.0	120
149	There's No Place Like Home: Crown-of-Thorns Outbreaks in the Central Pacific Are Regionally Derived and Independent Events. <i>PLoS ONE</i> , 2012, 7, e31159.	1.1	53
150	On the status of the Hawaiian seahorses <i>Hippocampus hilonis</i> , <i>H. histrix</i> and <i>H. fisheri</i> (Syngnathidae). <i>Marine Biology Research</i> , 2011, 7, 701-709.	0.3	5
151	New Records of Commercially Valuable Black Corals (Cnidaria: Antipatharia) from the Northwestern Hawaiian Islands at Mesophotic Depths. <i>Pacific Science</i> , 2011, 65, 249-255.	0.2	11
152	Coastal pollution limits pelagic larval dispersal. <i>Nature Communications</i> , 2011, 2, 226.	5.8	56
153	Marine connectivity: a new look at pelagic larval duration and genetic metrics of dispersal. <i>Marine Ecology - Progress Series</i> , 2011, 436, 291-305.	0.9	307
154	Discordant Phylogeographic and Biogeographic Breaks in California Halibut. <i>Bulletin (Southern)</i> 10 Tf 50 38	0.1	2
155	Widespread Dispersal of the Crown-of-Thorns Sea Star, <i>Acanthaster planci</i> , across the Hawaiian Archipelago and Johnston Atoll. <i>Journal of Marine Biology</i> , 2011, 2011, 1-10.	1.0	23
156	Defining Boundaries for Ecosystem-Based Management: A Multispecies Case Study of Marine Connectivity across the Hawaiian Archipelago. <i>Journal of Marine Biology</i> , 2011, 2011, 1-13.	1.0	116
157	Genetic Analyses and Simulations of Larval Dispersal Reveal Distinct Populations and Directional Connectivity across the Range of the Hawaiian Grouper (<i>Epinephelus quernus</i>). <i>Journal of Marine Biology</i> , 2011, 2011, 1-11.	1.0	23
158	Gateways to Hawaii: Genetic Population Structure of the Tropical Sea Cucumber <i>Holothuria atra</i> . <i>Journal of Marine Biology</i> , 2011, 2011, 1-16.	1.0	35
159	Variation in Symbiodinium ITS2 Sequence Assemblages among Coral Colonies. <i>PLoS ONE</i> , 2011, 6, e15854.	1.1	101
160	Detecting and measuring genetic differentiation. <i>Crustacean Issues</i> , 2011, , 31-55.	0.9	86
161	Phylogeography of <i>Emerita analoga</i> (Crustacea, Decapoda, Hippidae), an eastern Pacific Ocean sand crab with long-lived pelagic larvae. <i>Journal of Biogeography</i> , 2011, 38, 1600-1612.	1.4	34
162	Diversification of sympatric broadcast-spawning limpets (<i>Cellana</i> spp.) within the Hawaiian archipelago. <i>Molecular Ecology</i> , 2011, 20, 2128-2141.	2.0	79

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163	Range-Wide Genetic Connectivity of the Hawaiian Monk Seal and Implications for Translocation. <i>Conservation Biology</i> , 2011, 25, 124-132.	2.4	17
164	Sexual reproduction of Hawaiian black corals, with a review of the reproduction of antipatharians (Cnidaria: Anthozoa: Hexacorallia). <i>Invertebrate Biology</i> , 2011, 130, 211-225.	0.3	25
165	Preservation of corals in salt-saturated DMSO buffer is superior to ethanol for PCR experiments. <i>Coral Reefs</i> , 2011, 30, 329-333.	0.9	63
166	Azooxanthellate? Most Hawaiian black corals contain <i>Symbiodinium</i> . <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2011, 278, 1323-1328.	1.2	39
167	Inter-Specific Coral Chimerism: Genetically Distinct Multicellular Structures Associated with Tissue Loss in <i>Montipora capitata</i> . <i>PLoS ONE</i> , 2011, 6, e22869.	1.1	27
168	Escaping paradise: larval export from Hawaii in an Indo-Pacific reef fish, the yellow tang <i>Zebrasoma flavescens</i> . <i>Marine Ecology - Progress Series</i> , 2011, 428, 245-258.	0.9	55
169	Genetic evaluation of marine biogeographical barriers: perspectives from two widespread Indo-Pacific snappers (<i>Lutjanus kasmira</i> and <i>Lutjanus fulvus</i>). <i>Journal of Biogeography</i> , 2010, 37, 133-147.	1.4	161
170	Community ecology of mesophotic coral reef ecosystems. <i>Coral Reefs</i> , 2010, 29, 255-275.	0.9	386
171	Development of microsatellite markers from four Hawaiian corals: <i>Acropora cytherea</i> , <i>Fungia scutaria</i> , <i>Montipora capitata</i> and <i>Porites lobata</i> . <i>Conservation Genetics Resources</i> , 2010, 2, 11-15.	0.4	21
172	Isolation and characterization of microsatellite markers for the Crimson Jobfish, <i>Pristipomoides filamentosus</i> (Lutjanidae). <i>Conservation Genetics Resources</i> , 2010, 2, 169-172.	0.4	14
173	Rolling stones and stable homes: social structure, habitat diversity and population genetics of the Hawaiian spinner dolphin (<i>Stenella longirostris</i>). <i>Molecular Ecology</i> , 2010, 19, 732-748.	2.0	92
174	Genetic consequences of introducing allopatric lineages of Bluestriped Snapper (<i>Lutjanus kasmira</i>) to Hawaii. <i>Molecular Ecology</i> , 2010, 19, 1107-1121.	2.0	37
175	Protein expression and genetic structure of the coral <i>Porites lobata</i> in an environmentally extreme Samoan back reef: does host genotype limit phenotypic plasticity?. <i>Molecular Ecology</i> , 2010, 19, 1705-1720.	2.0	199
176	Taking the chaos out of genetic patchiness: seascape genetics reveals ecological and oceanographic drivers of genetic patterns in three temperate reef species. <i>Molecular Ecology</i> , 2010, 19, 3708-3726.	2.0	252
177	Isolation by distance across the Hawaiian Archipelago in the reef-building coral <i>Porites lobata</i> . <i>Molecular Ecology</i> , 2010, 19, 4661-4677.	2.0	54
178	Ocean currents help explain population genetic structure. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2010, 277, 1685-1694.	1.2	398
179	Is multiple mating beneficial or unavoidable? Low multiple paternity and genetic diversity in the shortspine spurdog <i>Squalus mitsukurii</i> . <i>Marine Ecology - Progress Series</i> , 2010, 403, 255-267.	0.9	63
180	Using morphometrics, in situ observations and genetic characters to distinguish among commercially valuable Hawaiian black coral species; a redescription of <i>Antipathes grandis</i> Verrill, 1928 (Antipatharia: Antipathidae). <i>Invertebrate Systematics</i> , 2010, 24, 271.	0.5	28

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181	Resurrection of <i>Porites hawaiiensis</i> Vaughan, 1907; a Hawaiian coral obscured by small size, cryptic habitat, and confused taxonomy. <i>Zootaxa</i> , 2010, 2624, .	0.2	6
182	Ecomorph or Endangered Coral? DNA and Microstructure Reveal Hawaiian Species Complexes: <i>Montipora dilatata/flabellata/turgescens</i> & <i>M. patula/verrilli</i> . <i>PLoS ONE</i> , 2010, 5, e15021.	1.1	56
183	Resolving natural ranges and marine invasions in a globally distributed octocoral (genus <i>Carijoa</i>). <i>Marine Ecology - Progress Series</i> , 2010, 401, 113-127.	0.9	55
184	Observations on the life history and feeding ecology of a specialized nudibranch predator (<i>Phyllodesmium poindimiei</i>), with implications for biocontrol of an invasive octocoral (<i>Carijoa riisei</i>) in Hawaii. <i>Journal of Experimental Marine Biology and Ecology</i> , 2009, 372, 64-74.	0.7	26
185	Shape-shifting corals: Molecular markers show morphology is evolutionarily plastic in <i>Porites</i> . <i>BMC Evolutionary Biology</i> , 2009, 9, 45.	3.2	183
186	Characterization of eight polymorphic microsatellite loci for the California spiny lobster, <i>Panulirus interruptus</i> and cross-amplification in other achelate lobsters. <i>Conservation Genetics Resources</i> , 2009, 1, 193-197.	0.4	10
187	A map of human impacts to a "pristine" coral reef ecosystem, the Papahānaumokuākea Marine National Monument. <i>Coral Reefs</i> , 2009, 28, 635-650.	0.9	111
188	Endemism and dispersal: comparative phylogeography of three surgeonfishes across the Hawaiian Archipelago. <i>Marine Biology</i> , 2009, 156, 689-698.	0.7	65
189	Generalist dinoflagellate endosymbionts and host genotype diversity detected from mesophotic (67-100 m depths) coral <i>Leptoseris</i> . <i>BMC Ecology</i> , 2009, 9, 21.	3.0	29
190	DISCORDANT DISTRIBUTION OF POPULATIONS AND GENETIC VARIATION IN A SEA STAR WITH HIGH DISPERSAL POTENTIAL. <i>Evolution; International Journal of Organic Evolution</i> , 2009, 63, 3214-3227.	1.1	55
191	Population genetics, larval dispersal, and connectivity in marine systems. <i>Marine Ecology - Progress Series</i> , 2009, 393, 1-12.	0.9	442
192	An alternative to ITS, a hypervariable, single-copy nuclear intron in corals, and its use in detecting cryptic species within the octocoral genus <i>Carijoa</i> . <i>Coral Reefs</i> , 2008, 27, 323-336.	0.9	52
193	Evaluating anthropogenic threats to the Northwestern Hawaiian Islands. <i>Aquatic Conservation: Marine and Freshwater Ecosystems</i> , 2008, 18, 1149-1165.	0.9	32
194	Isolation and characterization of polymorphic microsatellite and COI loci from the whelk <i>Kelletia kelletii</i> . <i>Molecular Ecology Resources</i> , 2008, 8, 881-883.	2.2	6
195	Extremely Low Genetic Diversity in the Endangered Hawaiian Monk Seal (<i>Monachus schauinslandi</i>). <i>Journal of Heredity</i> , 2008, 100, 25-33.	1.0	57
196	Isolation and characterization of polymorphic microsatellite and COI loci from the whelk <i>Kelletia kelletii</i> . <i>Molecular Ecology Resources</i> , 2008, .	2.2	0
197	An Evaluation of Cryptic Lineages of <i>Idotea Balthica</i> (Isopoda: Idoteidae): Morphology and Microsatellites. <i>Journal of Crustacean Biology</i> , 2007, 27, 643-648.	0.3	7
198	Host shift and speciation in a coral-feeding nudibranch. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2007, 274, 111-119.	1.2	76

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199	Frequency of multiple paternity in an unexploited tropical population of sandbar sharks (<i>Carcharhinus plumbeus</i>). <i>Canadian Journal of Fisheries and Aquatic Sciences</i> , 2007, 64, 198-204.	0.7	44
200	Contrasting phylogeography in three endemic Hawaiian limpets (<i>Cellana</i> spp.) with similar life histories. <i>Molecular Ecology</i> , 2007, 16, 3173-3186.	2.0	123
201	New report of nudibranch predators of the invasive octocoral <i>Carijoa riisei</i> in the Main Hawaiian Islands. <i>Coral Reefs</i> , 2007, 26, 411-411.	0.9	4
202	If larvae were smart: a simple model for optimal settlement behavior of competent larvae. <i>Marine Ecology - Progress Series</i> , 2007, 349, 43-61.	0.9	18
203	Spatial variability of recruitment in the sand crab <i>Emerita analoga</i> throughout California in relation to wind-driven currents. <i>Marine Ecology - Progress Series</i> , 2007, 350, 1-17.	0.9	27
204	Microsatellites for ecologists: a practical guide to using and evaluating microsatellite markers. <i>Ecology Letters</i> , 2006, 9, 615-629.	3.0	1,217
205	Noncoding mitochondrial loci for corals. <i>Molecular Ecology Notes</i> , 2006, 6, 1208-1211.	1.7	25
206	Assessment of multiple paternity in single litters from three species of carcharhinid sharks in Hawaii. <i>Environmental Biology of Fishes</i> , 2006, 76, 419-424.	0.4	46
207	Mitochondrial DNA and population size. <i>Science</i> , 2006, 314, 1388-90; author reply 1388-90.	6.0	3
208	Conservation implications of complex population structure: lessons from the loggerhead turtle (<i>Caretta caretta</i>). <i>Molecular Ecology</i> , 2005, 14, 2389-2402.	2.0	185
209	FOUNDATIONS OF GREGARIOUSNESS IN BARNACLES. <i>Journal of Experimental Biology</i> , 2005, 208, 1773-1774.	0.8	2
210	An experimental comparison of sediment-based biological filtration designs for recirculating aquarium systems. <i>Aquaculture</i> , 2005, 250, 244-255.	1.7	7
211	Genetic evidence of multiple paternity of broods in the intertidal crab <i>Petrolisthes cinctipes</i> . <i>Marine Ecology - Progress Series</i> , 2004, 270, 259-263.	0.9	47
212	Isolation and characterization of polymorphic microsatellite loci from the Dungeness crab <i>Cancer magister</i> . <i>Molecular Ecology Notes</i> , 2003, 4, 30-32.	1.7	11
213	FOUNDATIONS OF GREGARIOUSNESS: A DISPERSAL POLYMORPHISM AMONG THE PLANKTONIC LARVAE OF A MARINE INVERTEBRATE. <i>Evolution; International Journal of Organic Evolution</i> , 2001, 55, 2439.	1.1	17
214	FOUNDATIONS OF GREGARIOUSNESS: A DISPERSAL POLYMORPHISM AMONG THE PLANKTONIC LARVAE OF A MARINE INVERTEBRATE. <i>Evolution; International Journal of Organic Evolution</i> , 2001, 55, 2439-2454.	1.1	66
215	Settlement of the gregarious tube worm <i>Hydroides dianthus</i> (Polychaeta: Serpulidae). I. Gregarious and nongregarious settlement. <i>Marine Ecology - Progress Series</i> , 2001, 224, 103-114.	0.9	38
216	Settlement of the gregarious tube worm <i>Hydroides dianthus</i> (Polychaeta: Serpulidae). II. Testing the desperate larva hypothesis. <i>Marine Ecology - Progress Series</i> , 2001, 224, 115-131.	0.9	57

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217	Increased throughput for fragment analysis on an ABI PRISM 377 automated sequencer using a membrane comb and STRand software. <i>BioTechniques</i> , 2001, 31, 1320-4.	0.8	214
218	Settlement of the tube worm <i>Hydroides dianthus</i> (Polychaeta: Serpulidae): cues for gregarious settlement. <i>Marine Biology</i> , 1996, 126, 725-733.	0.7	66
219	Defenses of Caribbean sponges against predatory reef fish. I. Chemical detergency. <i>Marine Ecology - Progress Series</i> , 1995, 127, 183-194.	0.9	330
220	Foundations of gregariousness. <i>Nature</i> , 1994, 370, 511-512.	13.7	108
221	Limitations of laboratory assessments of coelenterate predation: Container effects on the prey selection of the <i>Limnomedusa</i> , <i>Proboscoidactyla Flaviccirrata</i> (Brandt). <i>Journal of Experimental Marine Biology and Ecology</i> , 1993, 167, 215-235.	0.7	26
222	Phylogenomics of <i>Palythoa</i> (Hexacorallia: Scleractinia): probing species boundaries in a globally distributed genus. <i>Coral Reefs</i> , 0, , 1.	0.9	2
223	Fish Flow: following fisheries from spawning to supper. <i>Frontiers in Ecology and the Environment</i> , 0, , .	1.9	3
224	Nitric oxide production rather than oxidative stress and cell death is associated with the onset of coral bleaching in <i>Pocillopora acuta</i> . <i>PeerJ</i> , 0, 10, e13321.	0.9	3
225	Coral micro-fragmentation assays for optimizing active reef restoration efforts. <i>PeerJ</i> , 0, 10, e13653.	0.9	4